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Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: [year=2010; month=5; day=13; hr=13; min=47; sec=54; ms=21;]

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Application No: 10580782 Version No: 2.0

Input Set:**Output Set:**

Started: 2010-05-10 14:32:26.808
Finished: 2010-05-10 14:32:29.464
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 656 ms
Total Warnings: 21
Total Errors: 0
No. of SeqIDs Defined: 21
Actual SeqID Count: 21

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (2)
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W 213	Artificial or Unknown found in <213> in SEQ ID (5)
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W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)

Input Set:

Output Set:

Started: 2010-05-10 14:32:26.808
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Total Warnings: 21
Total Errors: 0
No. of SeqIDs Defined: 21
Actual SeqID Count: 21

Error code

Error Description

This error has occurred more than 20 times, will not be displayed

SEQUENCE LISTING

<110> Miura, Yoshiko
Shibata, Chieri
Kobayashi, ChKazukiyo

<120> Glycopeptides and Temperature-Responsive Micelles

<130> TESHPl04US

<140> 10580782

<141> 2010-05-10

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<170> PatentIn version 3.5

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1 5

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<223> A sugar attached to C-term via a linker

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1 5 10

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1 5

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<223> 1 to 2 repeats for a minimum of 5 amino acid residues and a
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<221> CARBOHYD

<222> (5)..(5)

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linker

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1 5

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<221> MISC_FEATURE

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<223> Xaa is any amino acid

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<221> CARBOHYD

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 1 5

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<400> 9

Glu Val Pro Gly Val Gly
1 5

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1 5

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of 10 amino acid residues

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<223> Xaa is any amino acid residue

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1 5

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 linker

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<220>
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maximum of 50 amino acid residues

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